

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.

Application Serial Number: 10/590,579
Source: IFWP
Date Processed by STIC: 9/1/06

ENTERED



IFWP

RAW SEQUENCE LISTING

DATE: 09/01/2006

PATENT APPLICATION: US/10/590,579

TIME: 11:48:45

Input Set : A:\21491Y PCT.TXT

Output Set: N:\CRF4\09012006\J590579.raw

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4 <110> APPLICANT: Merck & Co., Inc.
6 <120> TITLE OF INVENTION: POLYPEPTIDES FOR INDUCING A PROTECTIVE
7   IMMUNE RESPONSE AGAINST STAPHYLOCOCCUS AUREUS
10 <130> FILE REFERENCE: 21491Y PCT
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/590,579
C--> 12 <141> CURRENT FILING DATE: 2006-08-22
12 <150> PRIOR APPLICATION NUMBER: 60/548,660
13 <151> PRIOR FILING DATE: 2004-02-27
15 <160> NUMBER OF SEQ ID NOS: 9
17 <170> SOFTWARE: FastSEQ for Windows Version 4.0
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 189
21 <212> TYPE: PRT
22 <213> ORGANISM: Artificial Sequence
24 <220> FEATURE:
25 <223> OTHER INFORMATION: truncated derivative of ORF0594
27 <400> SEQUENCE: 1
28 Arg Thr Asp Leu Lys Gly Ser Glu Phe Thr Phe Thr Pro Glu Ala Pro
29   1           5           10           15
30 Lys Thr Ile Thr Glu Leu Glu Lys Lys Val Glu Glu Ile Pro Phe Lys
31   20           25           30
32 Lys Glu Arg Lys Phe Asn Pro Asp Leu Ala Pro Gly Thr Glu Lys Val
33   35           40           45
34 Thr Arg Glu Gly Gln Lys Gly Glu Lys Thr Ile Thr Thr Pro Thr Leu
35   50           55           60
36 Lys Asn Pro Leu Thr Gly Val Ile Ile Ser Lys Gly Glu Pro Lys Glu
37 65           70           75           80
38 Glu Ile Thr Lys Asp Pro Ile Asn Glu Leu Thr Glu Tyr Gly Pro Glu
39   85           90           95
40 Thr Ile Ala Pro Gly His Arg Asp Glu Phe Asp Pro Lys Leu Pro Thr
41   100          105          110
42 Gly Glu Lys Glu Glu Val Pro Gly Lys Pro Gly Ile Lys Asn Pro Glu
43   115          120          125
44 Thr Gly Asp Val Val Arg Pro Pro Val Asp Ser Val Thr Lys Tyr Gly
45   130          135          140
46 Pro Val Lys Gly Asp Ser Ile Val Glu Lys Glu Glu Ile Pro Phe Glu
47 145          150          155          160
48 Lys Glu Arg Lys Phe Asn Pro Asp Leu Ala Pro Gly Thr Glu Lys Val
49   165          170          175
50 Thr Arg Glu Gly Gln Lys Gly Glu Lys Thr Ile Thr Thr
51   180          185
54 <210> SEQ ID NO: 2
55 <211> LENGTH: 235

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56 <212> TYPE: PRT

57 <213> ORGANISM: Artificial Sequence

59 <220> FEATURE:

60 <223> OTHER INFORMATION: His-tagged derivative of SEQ ID NO: 1

62 <400> SEQUENCE: 2

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63 Met His His His His His His Ser Ser Gly Leu Val Pro Arg Gly Ser
64 1          5          10          15
65 Gly Met Lys Glu Thr Ala Ala Ala Lys Phe Glu Arg Gln His Met Asp
66          20          25          30
67 Ser Pro Asp Leu Gly Thr Asp Asp Asp Asp Lys Ala Met Gly Arg Thr
68          35          40          45
69 Asp Leu Lys Gly Ser Glu Phe Thr Phe Thr Pro Glu Ala Pro Lys Thr
70          50          55          60
71 Ile Thr Glu Leu Glu Lys Lys Val Glu Glu Ile Pro Phe Lys Lys Glu
72 65          70          75          80
73 Arg Lys Phe Asn Pro Asp Leu Ala Pro Gly Thr Glu Lys Val Thr Arg
74          85          90          95
75 Glu Gly Gln Lys Gly Glu Lys Thr Ile Thr Thr Pro Thr Leu Lys Asn
76          100          105          110
77 Pro Leu Thr Gly Val Ile Ile Ser Lys Gly Glu Pro Lys Glu Glu Ile
78          115          120          125
79 Thr Lys Asp Pro Ile Asn Glu Leu Thr Glu Tyr Gly Pro Glu Thr Ile
80          130          135          140
81 Ala Pro Gly His Arg Asp Glu Phe Asp Pro Lys Leu Pro Thr Gly Glu
82 145          150          155          160
83 Lys Glu Glu Val Pro Gly Lys Pro Gly Ile Lys Asn Pro Glu Thr Gly
84          165          170          175
85 Asp Val Val Arg Pro Pro Val Asp Ser Val Thr Lys Tyr Gly Pro Val
86          180          185          190
87 Lys Gly Asp Ser Ile Val Glu Lys Glu Glu Ile Pro Phe Glu Lys Glu
88          195          200          205
89 Arg Lys Phe Asn Pro Asp Leu Ala Pro Gly Thr Glu Lys Val Thr Arg
90          210          215          220
91 Glu Gly Gln Lys Gly Glu Lys Thr Ile Thr Thr
92 225          230          235

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95 <210> SEQ ID NO: 3

96 <211> LENGTH: 1240

97 <212> TYPE: PRT

98 <213> ORGANISM: S. aureus

100 <400> SEQUENCE: 3

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101 Met Arg Asp Lys Lys Gly Pro Val Asn Lys Arg Val Asp Phe Leu Ser
102 1          5          10          15
103 Asn Lys Leu Asn Lys Tyr Ser Ile Arg Lys Phe Thr Val Gly Thr Ala
104          20          25          30
105 Ser Ile Leu Ile Gly Ser Leu Met Tyr Leu Gly Thr Gln Gln Glu Ala
106          35          40          45
107 Glu Ala Ala Glu Asn Asn Ile Glu Asn Pro Thr Thr Leu Lys Asp Asn
108          50          55          60
109 Val Gln Ser Lys Glu Val Lys Ile Glu Glu Val Thr Asn Lys Asp Thr

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110	65				70				75				80			
111	Ala	Pro	Gln	Gly	Val	Glu	Ala	Lys	Ser	Glu	Val	Thr	Ser	Asn	Lys	Asp
112					85					90					95	
113	Thr	Ile	Glu	His	Glu	Pro	Ser	Val	Lys	Ala	Glu	Asp	Ile	Ser	Lys	Lys
114					100					105					110	
115	Glu	Asp	Thr	Pro	Lys	Glu	Val	Ala	Asp	Val	Ala	Glu	Val	Gln	Pro	Lys
116					115					120					125	
117	Ser	Ser	Val	Thr	His	Asn	Ala	Glu	Thr	Pro	Lys	Val	Arg	Lys	Ala	Arg
118					130					135					140	
119	Ser	Val	Asp	Glu	Gly	Ser	Phe	Asp	Ile	Thr	Arg	Asp	Ser	Lys	Asn	Val
120	145					150					155					160
121	Val	Glu	Ser	Thr	Pro	Ile	Thr	Ile	Gln	Gly	Lys	Glu	His	Phe	Glu	Gly
122						165					170					175
123	Tyr	Gly	Ser	Val	Asp	Ile	Gln	Lys	Lys	Pro	Thr	Asp	Leu	Gly	Val	Ser
124					180					185					190	
125	Glu	Val	Thr	Arg	Phe	Asn	Val	Gly	Asn	Glu	Ser	Asn	Gly	Leu	Ile	Gly
126					195					200					205	
127	Ala	Leu	Gln	Leu	Lys	Asn	Lys	Ile	Asp	Phe	Ser	Lys	Asp	Phe	Asn	Phe
128					210					215					220	
129	Lys	Val	Arg	Val	Ala	Asn	Asn	His	Gln	Ser	Asn	Thr	Thr	Gly	Ala	Asp
130	225					230					235					240
131	Gly	Trp	Gly	Phe	Leu	Phe	Ser	Lys	Gly	Asn	Ala	Glu	Glu	Tyr	Leu	Thr
132					245					250					255	
133	Asn	Gly	Gly	Ile	Leu	Gly	Asp	Lys	Gly	Leu	Val	Asn	Ser	Gly	Gly	Phe
134					260					265					270	
135	Lys	Ile	Asp	Thr	Gly	Tyr	Ile	Tyr	Thr	Ser	Ser	Met	Asp	Lys	Thr	Glu
136					275					280					285	
137	Lys	Gln	Ala	Gly	Gln	Gly	Tyr	Arg	Gly	Tyr	Gly	Ala	Phe	Val	Lys	Asn
138					290					295					300	
139	Asp	Ser	Ser	Gly	Asn	Ser	Gln	Met	Val	Gly	Glu	Asn	Ile	Asp	Lys	Ser
140	305					310					315				320	
141	Lys	Thr	Asn	Phe	Leu	Asn	Tyr	Ala	Asp	Asn	Ser	Thr	Asn	Thr	Ser	Asp
142					325					330					335	
143	Gly	Lys	Phe	His	Gly	Gln	Arg	Leu	Asn	Asp	Val	Ile	Leu	Thr	Tyr	Val
144					340					345					350	
145	Ala	Ser	Thr	Gly	Lys	Met	Arg	Ala	Glu	Tyr	Ala	Gly	Lys	Thr	Trp	Glu
146					355					360					365	
147	Thr	Ser	Ile	Thr	Asp	Leu	Gly	Leu	Ser	Lys	Asn	Gln	Ala	Tyr	Asn	Phe
148					370					375					380	
149	Leu	Ile	Thr	Ser	Ser	Gln	Arg	Trp	Gly	Leu	Asn	Gln	Gly	Ile	Asn	Ala
150	385					390					395					400
151	Asn	Gly	Trp	Met	Arg	Thr	Asp	Leu	Lys	Gly	Ser	Glu	Phe	Thr	Phe	Thr
152					405					410					415	
153	Pro	Glu	Ala	Pro	Lys	Thr	Ile	Thr	Glu	Leu	Glu	Lys	Lys	Val	Glu	Glu
154					420					425					430	
155	Ile	Pro	Phe	Lys	Lys	Glu	Arg	Lys	Phe	Asn	Pro	Asp	Leu	Ala	Pro	Gly
156					435					440					445	
157	Thr	Glu	Lys	Val	Thr	Arg	Glu	Gly	Gln	Lys	Gly	Glu	Lys	Thr	Ile	Thr
158					450					455					460	

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159 Thr Pro Thr Leu Lys Asn Pro Leu Thr Gly Val Ile Ile Ser Lys Gly
160 465 470 475 480
161 Glu Pro Lys Glu Glu Ile Thr Lys Asp Pro Ile Asn Glu Leu Thr Glu
162 485 490 495
163 Tyr Gly Pro Glu Thr Ile Ala Pro Gly His Arg Asp Glu Phe Asp Pro
164 500 505 510
165 Lys Leu Pro Thr Gly Glu Lys Glu Glu Val Pro Gly Lys Pro Gly Ile
166 515 520 525
167 Lys Asn Pro Glu Thr Gly Asp Val Val Arg Pro Pro Val Asp Ser Val
168 530 535 540
169 Thr Lys Tyr Gly Pro Val Lys Gly Asp Ser Ile Val Glu Lys Glu Glu
170 545 550 555 560
171 Ile Pro Phe Glu Lys Glu Arg Lys Phe Asn Pro Asp Leu Ala Pro Gly
172 565 570 575
173 Thr Glu Lys Val Thr Arg Glu Gly Gln Lys Gly Glu Lys Thr Ile Thr
174 580 585 590
175 Thr Pro Thr Leu Lys Asn Pro Leu Thr Gly Val Ile Ile Ser Lys Gly
176 595 600 605
177 Glu Pro Lys Glu Glu Ile Thr Lys Asp Pro Ile Asn Glu Leu Thr Glu
178 610 615 620
179 Tyr Gly Pro Glu Thr Ile Thr Pro Gly His Arg Asp Glu Phe Asp Pro
180 625 630 635 640
181 Lys Leu Pro Thr Gly Glu Lys Glu Glu Val Pro Gly Lys Pro Gly Ile
182 645 650 655
183 Lys Asn Pro Glu Thr Gly Asp Val Val Arg Pro Pro Val Asp Ser Val
184 660 665 670
185 Thr Lys Tyr Gly Pro Val Lys Gly Asp Ser Ile Val Glu Lys Glu Glu
186 675 680 685
187 Ile Pro Phe Lys Lys Glu Arg Lys Phe Asn Pro Asp Leu Ala Pro Gly
188 690 695 700
189 Thr Glu Lys Val Thr Arg Glu Gly Gln Lys Gly Glu Lys Thr Ile Thr
190 705 710 715 720
191 Thr Pro Thr Leu Lys Asn Pro Leu Thr Gly Glu Ile Ile Ser Lys Gly
192 725 730 735
193 Glu Ser Lys Glu Glu Ile Thr Lys Asp Pro Ile Asn Glu Leu Thr Glu
194 740 745 750
195 Tyr Gly Pro Glu Thr Ile Thr Pro Gly His Arg Asp Glu Phe Asp Pro
196 755 760 765
197 Lys Leu Pro Thr Gly Glu Lys Glu Glu Val Pro Gly Lys Pro Gly Ile
198 770 775 780
199 Lys Asn Pro Glu Thr Gly Asp Val Val Arg Pro Pro Val Asp Ser Val
200 785 790 795 800
201 Thr Lys Tyr Gly Pro Val Lys Gly Asp Ser Ile Val Glu Lys Glu Glu
202 805 810 815
203 Ile Pro Phe Lys Lys Glu Arg Lys Phe Asn Pro Asp Leu Ala Pro Gly
204 820 825 830
205 Thr Glu Lys Val Thr Arg Glu Gly Gln Lys Gly Glu Lys Thr Ile Thr
206 835 840 845
207 Thr Pro Thr Leu Lys Asn Pro Leu Thr Gly Glu Ile Ile Ser Lys Gly

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208      850      855      860
209 Glu Ser Lys Glu Glu Ile Thr Lys Asp Pro Ile Asn Glu Leu Thr Glu
210 865      870      875      880
211 Tyr Gly Pro Glu Thr Ile Thr Pro Gly His Arg Asp Glu Phe Asp Pro
212      885      890      895
213 Lys Leu Pro Thr Gly Glu Lys Glu Glu Val Pro Gly Lys Pro Gly Ile
214      900      905      910
215 Lys Asn Pro Glu Thr Gly Asp Val Val Arg Pro Pro Val Asp Ser Val
216      915      920      925
217 Thr Lys Tyr Gly Pro Val Lys Gly Asp Ser Ile Val Glu Lys Glu Glu
218      930      935      940
219 Ile Pro Phe Glu Lys Glu Arg Lys Phe Asn Pro Asp Leu Ala Pro Gly
220 945      950      955      960
221 Thr Glu Lys Val Thr Arg Glu Gly Gln Lys Gly Glu Lys Thr Ile Thr
222      965      970      975
223 Thr Pro Thr Leu Lys Asn Pro Leu Thr Gly Glu Ile Ile Ser Lys Gly
224      980      985      990
225 Glu Ser Lys Glu Glu Ile Thr Lys Asp Pro Val Asn Glu Leu Thr Glu
226      995      1000      1005
227 Phe Gly Gly Glu Lys Ile Pro Gln Gly His Lys Asp Ile Phe Asp Pro
228      1010      1015      1020
229 Asn Leu Pro Thr Asp Gln Thr Glu Lys Val Pro Gly Lys Pro Gly Ile
230 1025      1030      1035      1040
231 Lys Asn Pro Asp Thr Gly Lys Val Ile Glu Glu Pro Val Asp Asp Val
232      1045      1050      1055
233 Ile Lys His Gly Pro Lys Thr Gly Thr Pro Glu Thr Lys Thr Val Glu
234      1060      1065      1070
235 Ile Pro Phe Glu Thr Lys Arg Glu Phe Asn Pro Lys Leu Gln Pro Gly
236      1075      1080      1085
237 Glu Glu Arg Val Lys Gln Glu Gly Gln Pro Gly Ser Lys Thr Ile Thr
238      1090      1095      1100
239 Thr Pro Ile Thr Val Asn Pro Leu Thr Gly Glu Lys Val Gly Glu Gly
240 1105      1110      1115      1120
241 Gln Pro Thr Glu Glu Ile Thr Lys Gln Pro Val Asp Lys Ile Val Glu
242      1125      1130      1135
243 Phe Gly Gly Glu Lys Pro Lys Asp Pro Lys Gly Pro Glu Asn Pro Glu
244      1140      1145      1150
245 Lys Pro Ser Arg Pro Thr His Pro Ser Gly Pro Val Asn Pro Asn Asn
246      1155      1160      1165
247 Pro Gly Leu Ser Lys Asp Arg Ala Lys Pro Asn Gly Pro Val His Ser
248      1170      1175      1180
249 Met Asp Lys Asn Asp Lys Val Lys Lys Ser Lys Ile Ala Lys Glu Ser
250 1185      1190      1195      1200
251 Val Ala Asn Gln Glu Lys Lys Arg Ala Glu Leu Pro Lys Thr Gly Leu
252      1205      1210      1215
253 Glu Ser Thr Gln Lys Gly Leu Ile Phe Ser Ser Ile Ile Gly Ile Ala
254      1220      1225      1230
255 Gly Leu Met Leu Leu Ala Arg Arg
256      1235      1240

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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/590,579

DATE: 09/01/2006

TIME: 11:48:46

Input Set : A:\21491Y PCT.TXT

Output Set: N:\CRF4\09012006\J590579.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date